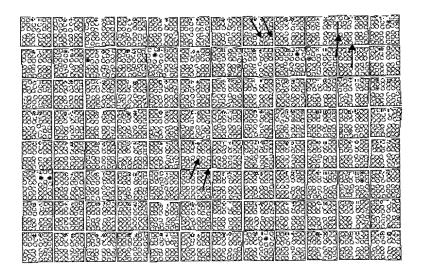
Figure 1

 \mathbf{A}



В

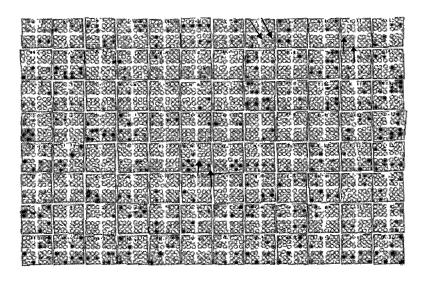


Figure 2

Differentially expressed genes	intact skin	wound	intact skin	wound
	control animals	control animals	dexamethasone	dexamethasone
SW1368	1,00	148,90	0,70	181,50
SW1695	1,00	0,24	0,94	0,61
Differentially expressed genes	intact skin	wound	intact skin	wound
	young mice	young mice	old mice	old mice
SW1368	1,20	170,70	0,80	204,60
SW1695	0,44	0,07	0,65	0,10

Figure 3

Differentially expressed genes	intact skin	wound	intact skin	wound
	control mice	control mice	diabetic mice	diabetic mice
SW1368	1,00	561,60	2,20	284,40
SW1695	1,00	0,07	1,01	0,08

Figure 4

CDNA-HUMAN*	SEQ ID No. 6	SEQ ID No. 8
CDNA-MOUSE*	SEQ ID No. 5	SEQ ID No. 7
Seq ID No.	7	4
-MOUSE* Seq ID No. PROTEIN-HUMAN* Seq ID No. CDNA-MOUSE* CDNA-HUMAN*	SEQ ID No. 2	SEQ ID No. 4
Seq ID No.	_	3
No. NAME PROTEIN-MOUSE*	SEO ID No. 1	SEQ ID No. 3
, NAME	SW1368	1. SW1695 2. SW1695
ž		. 2

^{*:} SEQ ID numbers of the sequence listing

Figure 5

SGT ---- VESALNYSRGSTVHTAYLVLSSLAMFTCLCGMAGNSMV Hum_sw1368 SASPTAPVTTNPMDETLPGSINIRILIPKLMIIIFGLVGLMGNAIV Mouse_sw1368 MGESNGEAFLAFKT -INS 1 MNQT

--IFHLCFNTIM Mouse_sw1368 Hum_sw1368 49 IWLLGFRMHRNPFCIYILNLAAADLLFLFS - - MASTLSLETQPLVNTTDKVHEL - MKRLM 61 FWLLGEHLRRNAESVYILNLALADFLFLLSSITASTLFLLKVSYLSI - - - IFHLCFNTIM

106YFAYTVGLSLLTAISTORCLSVLFPIWFKCHRPRHLSAWVCGLLWTLCLLMNGLTSSFCS Hum_sw1368 118MVVYITGISMLSAISTECCLSVLCBTWYRCHRBVHTSTVMCAVIWVLSLLICILNSYFGA Mouse_sw1368

166KFI - KFNEDR - CFRVDMVQAALIMGVLTPVMTLSSLTLFVWVRRSSQQWRRQPTRLFVVV Hum_sw1368 178VLHTRYDNDNEGLATNIFTASYMIFLLV-VLCLSSLALLARLFCGAGG--MKLTRFHYTI Mouse_sw1368

224LASVLVFLICSLPLSIYWFVLYWLSLPPEMQVL ---CFSLSRLSSSVSSSANPVIYFLV Hum_sw1368 2351LTTLVFLLGGLPFVLYCILLF--KIKDDFHVLDVNLYLALEVLTA-INSCANPILYEFV Mouse_sw1368

Mouse_sw1368 Hum_sw1368 280GSRRSHRLPTRSLGTVLQQALREEPE -----LEGGETPTVGTNEMG----A 292GSFR-HQLKHQTLKMVLQSALQDTPETAENMVEMSSNKAEPGGGHVYPYDVPDYA

All boxed mouse sequences exactly match the human sequence Hum_sw1368.

Figure 6

MDPTTPAWGTESTTVNGNDQALLLLCGKETLIPVFLILFIALVGLVGNGFVLWLLGFRMR Hum_sw1695 MDIDISSLGIYIIAPNGSSYTNSVDQFFKIQVMGFLSLILISPVGMVLNSTVLWFLGFQIR Mouse_sw1695

FICEQIINCLVYLSNFFCSISINFPSFFTTVMTCAYLAGLSMI Hum_sw1695 FIHSQF---LFYILAIFPSIPIQIEIDMLTKFAYISGLSII Mouse_sw1695 61 RNAFSVYVLSLAGADFL 61 RNAFSVYILNLAGADFL

STVSTERCLSVLWPIWYRCRRPRHLSAVVCVLLWALSLLLSILEGKFCGFLFSDGDSGWO Hum_sw1695 STISTERCIOVMWPIWYROORPRHTSSVTOSLLWALSLLFALLDGMGCGLLFNSFDQSWO Mouse_sw1695

18 OTFDFITAAWLIFLFMVLCGSSLALIJ-VRILCGSRGLPLTRLYLTILLTVLVFLLCGLPF Hum_sw1695 17 LKFDLJICAWSJVLFVVLCGSSJHPTCXRJFCGSQQIBVTRLYVTJALTVJSFLJGGLPF Mouse_sw1695

GIQWFLILWIWKDSDVLFCHIHPVSVVLSSLNSSANPIIYFFVGSFRKQWRLQQPILKLA Hum_sw1695 GISW-IIQWSETLIYVGFGDYFHEELFLSCINSCANPIIYFIVG-FIRGRKFQGKSLKVL Mouse_sw1695

LORALODIAEVDHSEGCFROGTPEMSRSSLV LORAMEDTPEENEDMGPSRNPEEFETVCSN 30

Hum_sw1695 Mouse_sw1695

> sequence Hum sw1695. All boxed mouse sequences exactly match the human

Figure 7

				Jose & wording	Dunon-14 wound
	into a frin	1 h wound	day-1 wound	day-5 wound	
	Illiact smill		1 07	1 50	5.21
Rel human SW1368 1,0	1,00	1,79	1,82	1,0,1	
Fxpression					

Figure 8

wound ground of ulcer patients	1 74	197	0,00	
wound edge of ulcer patients		0,00	00 0	0060
loow	Intact skill ulcei	100	2001	1,00
	Target gene	1 cW/1269	numan 5 w 1.300	human SW1695